

FIG. 1A



FIG. 1B

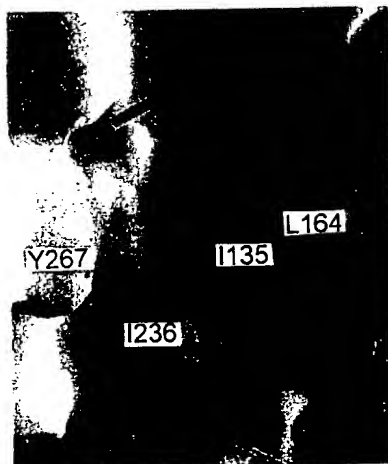


FIG. 1C



FIG. 1D

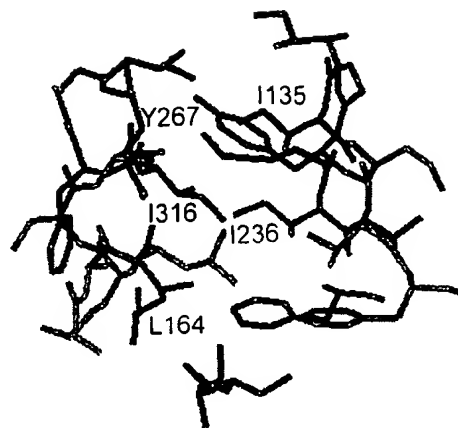
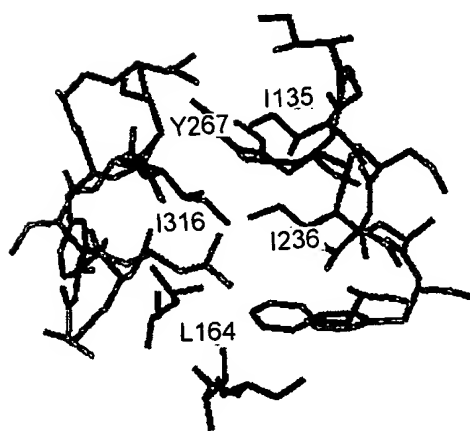


FIG. 1E

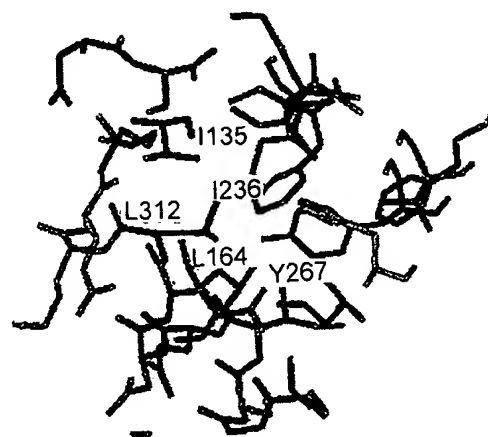
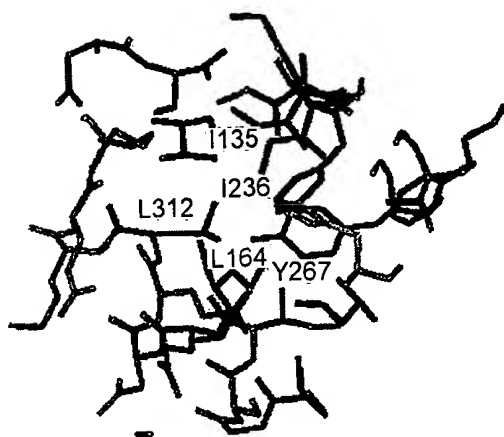


FIG. 1F

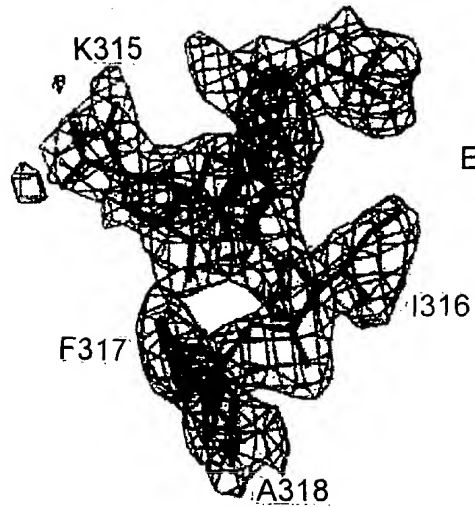


FIG. 2A

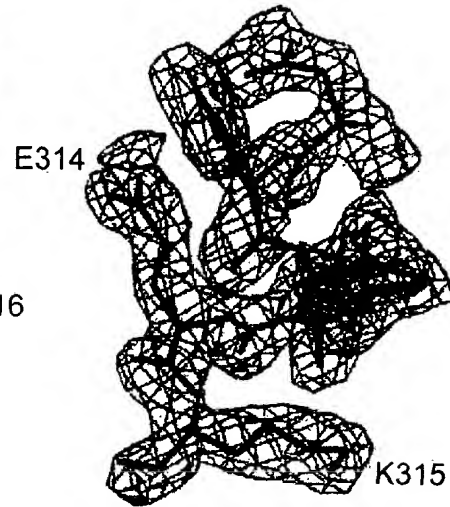


FIG. 2B



FIG. 2C

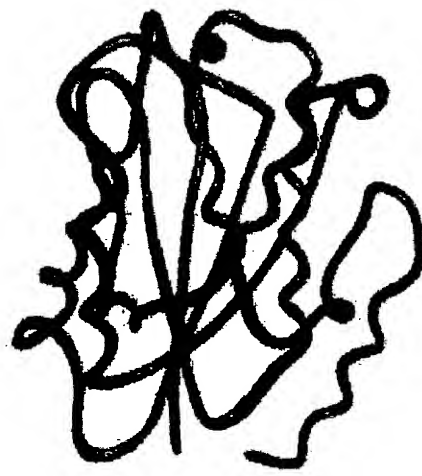


FIG. 2D

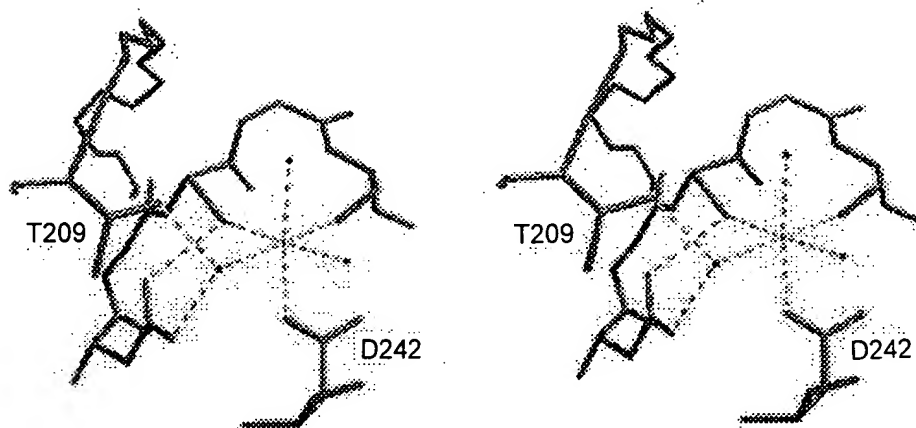


FIG. 2E

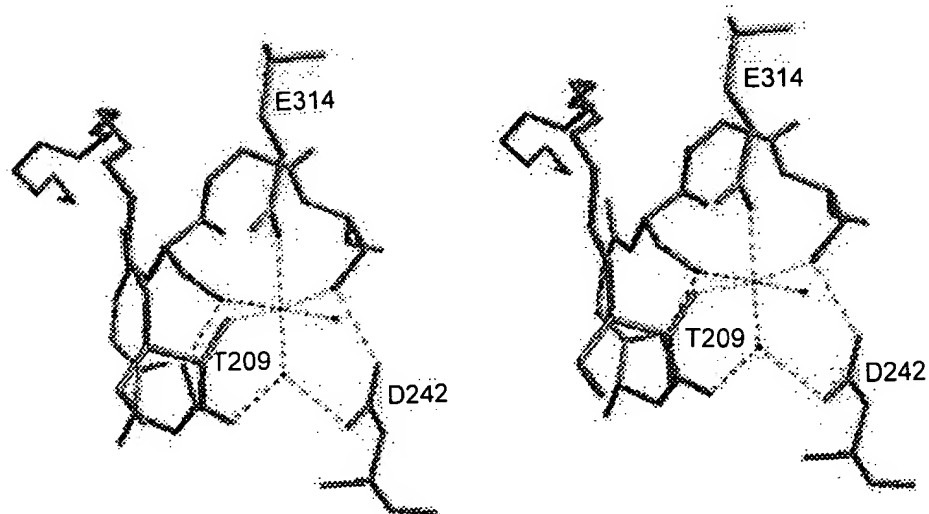
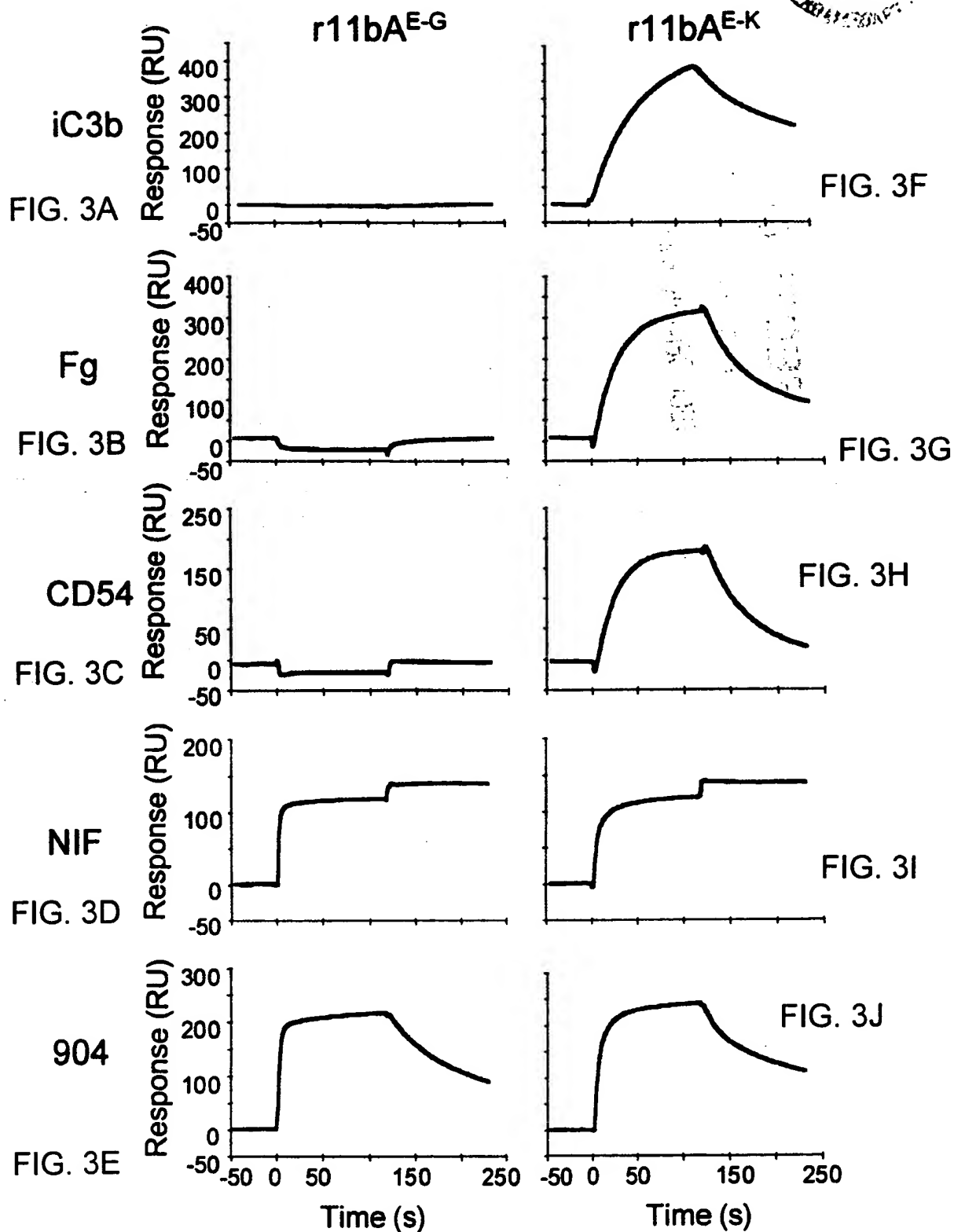
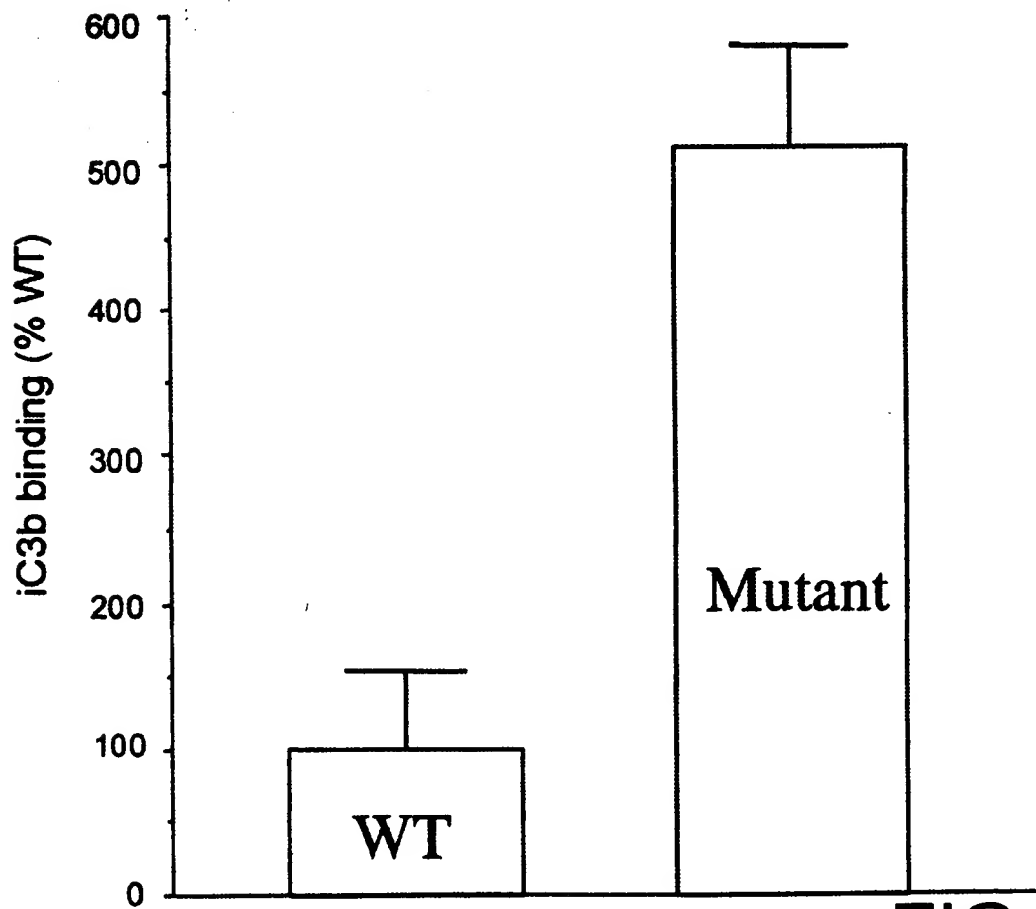
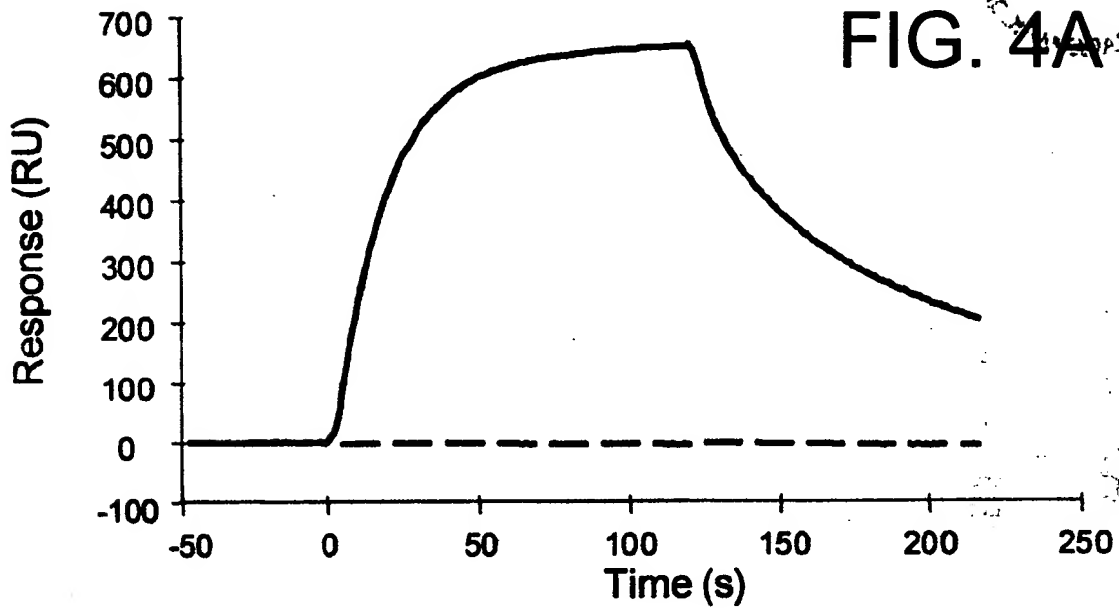


FIG. 2F





**FIG. 4B**

[illegible][illegible]

## consensus

11b-A 193.32 6:194  
11c-A 193.32 6:194  
11d-A 193.32 6:194  
11a-A 186.32 6:187  
a1-A 197.32 6:198  
a2-A 197.32 6:198  
a10-A 197.32 6:198  
a11-A 195.32 6:196  
a8-A 194.32 6:195  
consensus 201.1 \* :202

## consensus

FIG. 5

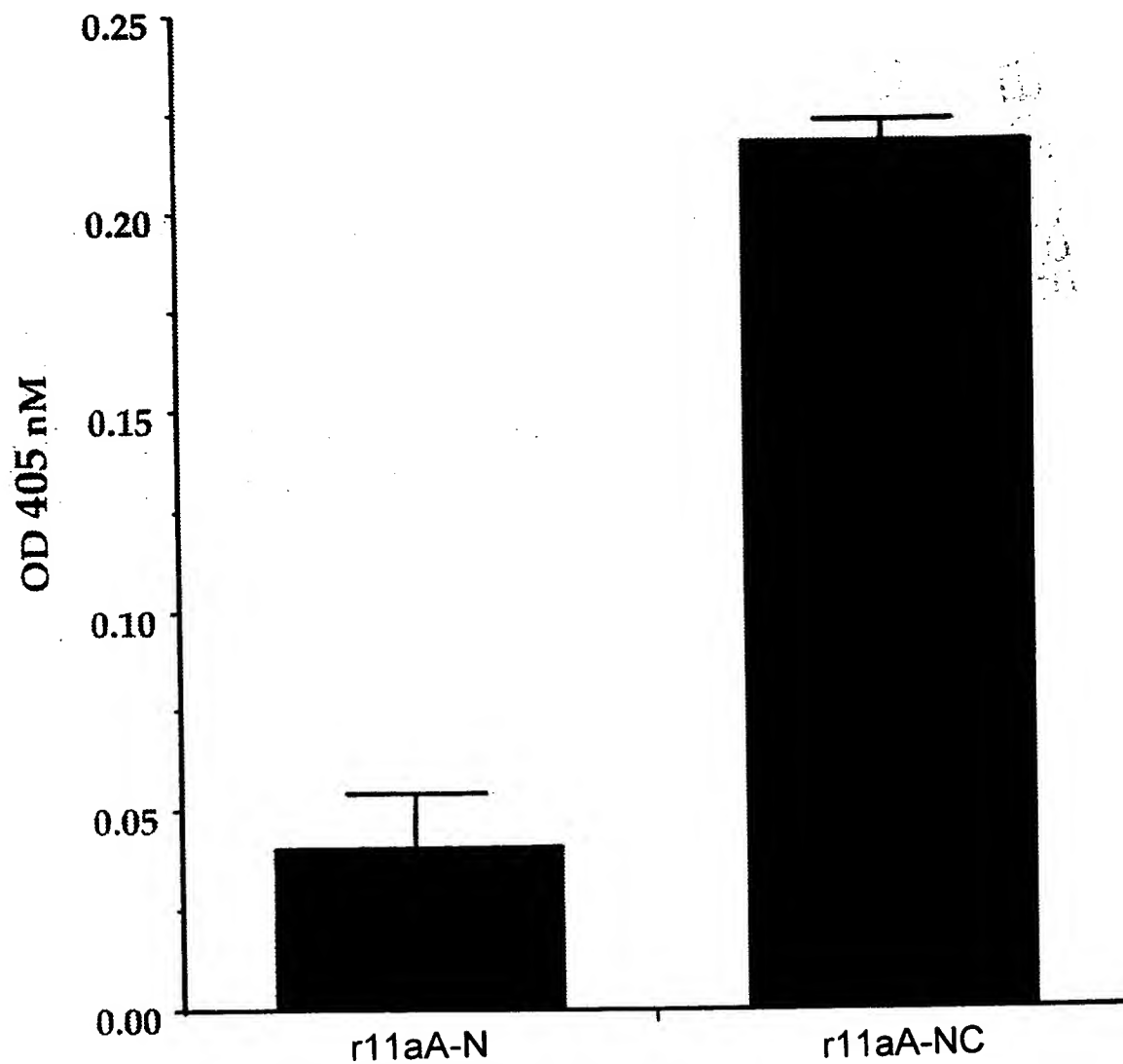


FIG. 6



# HIGH AFFINITY INTEGRIN POLYPEPTIDES AND USES THEREOF

```

170 180 190 200
Z Y R S S M S Q M S T K A T O R K R K
Z L A S S I A N S T K A E R K R K
Z L A D E S T A E S G S E S K R K
Z Y R O D S N O S T D M E R K R K
Z Y L O D R N O S G D L A N E S E
Z Y S L A R V O S H A L V A Q S H
Z Y A S S I O N S O N D S R S A P T
Z F N S S I O N S O N A R V S O S
I * * * * *

```

b3  
b5  
b6  
b1  
b2  
b7  
b8  
b4  
CO

[illegible]

b3  
b5  
b6  
b1  
b2  
b7  
b8  
b4  
conspays

[illegible]

b3  
b5  
b6  
b1  
b2  
b7  
b8  
b4  
consensus

410  
3370: TGG TGGKIR  
3371: TNN TSSIR  
3364: TIS TETR  
3371: TIG TSSS  
3356: TKN TTKS  
3382: TMD TTSB  
3377: TGG TTKI  
3359: TEG TTRR  
401: \*\*\*! \*\*--! \*

b3  
b5  
b6  
b1  
b2  
b7  
b8  
b4  
consensus

FIG. 7